

Bex Sequence

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CHEMOKINE AND RECEPTOR USES; COMPOSITIONS; METHODS

## Enclosed:

1. Request for a one-month extension of time (1 page, in duplicate); 2. Response to Notice to File Missing Parts (2 pages in duplicate); 3. Signed Declaration and Power of Attorney (Two copies signed in counterpart, 5 pages each); 4. Copy of Notice to File Missing Parts -Filing Date Granted (2 pages); 5. Response to Notice to Comply with Sequence Listing Rules (2 pages) 6. One write-protected diskette (CRM); 7. Paper print-out of contents of diskette (16 pages) 8. Return Postcard.

## SEQUENCE LISTING

<110> Oldham, Elizabeth K. Soto, Hortensia Liu, Ying Hudak, Susan A. Homey, Bernhard Morales, Janine M. Kellerman, Sirid-Aimee McEvoy, Leslie M. Bowman, Edward P. Zlotnik, Albert CHEMOKINE AND RECEPTOR USES; COMPOSITIONS; METHODS <130> DX0882XK US09/898,751 <140> <141> 2001-07-02 US09/471,549 <150> <151> 1999-12-23 <150> US60/136,570 <151> 1999-05-27 <150> US60/113,858 <151> 1998-12-24 <160> 16 <170> PatentIn version 3.1 <210> 1 <211> 1089 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (1)..(1086) <223> <400> 1 atg ggg acg gag gtt tta gag cag gtt tcc tgg ggc cat tac tct ggg 48 Met Gly Thr Glu Val Leu Glu Gln Val Ser Trp Gly His Tyr Ser Gly gat gaa gag gac gca tac tcg gct gag cca ctg ccg gag ctt tgc tac 96 Asp Glu Glu Asp Ala Tyr Ser Ala Glu Pro Leu Pro Glu Leu Cys Tyr 25 aag gcc gat gtc cag gcc ttc agc cgg gcc ttc caa ccc agt gtc tcc 144 Lys Ala Asp Val Gln Ala Phe Ser Arg Ala Phe Gln Pro Ser Val Ser 35 ctg acg ctg gct gcg ctg ggt ctg gcc ggc aat ggc ctg gtc ctg gcc 192 Leu Thr Leu Ala Ala Leu Gly Leu Ala Gly Asn Gly Leu Val Leu Ala

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	gcg Ala															336
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gat Asp	gtc Val 290	gca Ala	ctg Leu	ctg Leu	gtg Val	acc Thr 295	agc Ser	ggc Gly	ttg Leu	gcc Ala	ctc Leu 300	gcc Ala	cgc Arg	tgt Cys	ggc Gly	912
ctc	aat	ccc	gtt	ctc	tac	gcc	ttc	ctg	ggc	ctg	cgc	ttc	cgc	cag	gac	960

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Phe Ala Ala A	ala Gly Ala .00		Gly Trp Ser 105	Leu Gly Ser		
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ctc Leu	ttc Phe 130	cta Leu	gcc Ala	tgt Cys	atc Ile	agc Ser 135	gcc Ala	gac Asp	cgc Arg	tat Tyr	gtg Val 140	gcc Ala	atc Ile	gca Ala	cga Arg	432
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Thr His Leu Ala Ala Arg Arg Thr Thr Arg Ser Pro Thr Ser Val His

Leu Met Val Ala Val Leu Gly Leu Ala Gly Asn Gly Leu Val Leu Ala

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Leu Leu Gln Leu Ala Leu Ala Asp Leu Leu Leu Ala Leu Thr Leu Pro 85 90 95

Phe Ala Ala Ala Gly Ala Leu Gln Gly Trp Asn Leu Gly Ser Thr Thr
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Cys Arg Ala Ile Ser Gly Leu Tyr Ser Ala Ser Phe His Ala Gly Phe 115 120 125

Leu Phe Leu Ala Cys Ile Ser Ala Asp Arg Tyr Val Ala Ile Ala Arg 130 135 140

Ala Leu Pro Ala Gly Gln Arg Pro Ser Thr Pro Ser Arg Ala His Leu 145 150 155 160

Val Ser Val Phe Val Trp Leu Leu Ala Leu Phe Leu Ala Leu Pro Ala 165 170 175

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Val Ala Gln Val Val Leu Gly Phe Ala Leu Pro Leu Gly Val Met Ala 210 215 220

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Val Val Leu Gln Leu Pro Tyr Ser Leu Ala Leu Leu Leu Asp Thr Ala 260 265 270

Asp Leu Leu Ala Ala Arg Glu Arg Ser Cys Ser Ser Ser Lys Arg Lys 275 280 285

Asp Leu Ala Leu Leu Val Thr Gly Gly Leu Thr Leu Val Arg Cys Ser 290 295 300

Leu Asn Pro Val Leu Tyr Ala Phe Leu Gly Leu Arg Phe Arg Arg Asp 305 310 315 320

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His Val Lys Arg Arg Ile Cys Val Ser Pro His Asn His Thr Val

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taaa	atcto	cca a	agaat	gcca	at tt	ccct	atco	c cta	aatga	attc	aato	ctcc	ett a	accct	gacca	•	516
atca	agtgg	gee o	caaat	tttc	c ag	gccc	ettgo	e cto	ccag	gaac	ccca	gcc	cag a	aacto	cttcag	6	576
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Glu	Val	Ser	His	His 15	Ile	Ser	Arg	Arg	Leu 20	Leu	Glu	Arg	Val	Asn 25	Met		
Cys	Arg	Ile	Gln 30	Arg	Ala	Asp	Gly	Asp 35	Cys	Asp	Leu	Ala	Ala 40	Val	Ile		
Leu	His	Val 45	Lys	Arg	Arg	Arg	Ile 50	Cys	Val	Ser	Pro	His 55	Asn	His	Thr		
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Ala Pro Ser Leu Pro Leu Arg Ser Trp His Pro Trp Asn Lys Thr Lys 55 50

Gln Lys Gln Glu Ala Leu Pro Leu Pro Ser Ser Thr Ser Cys Cys Thr 70

Gln Leu Tyr Arg Gln Pro Leu Pro Ser Arg Leu Leu Arg Arg Ile Val

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Gln Ala Phe Val Leu His Leu Ala Gln Arg Ser Ile Cys Ile His Pro 45 50 55	
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Ser Cys Cys Thr Gln Leu Tyr Arg Gln Pro Leu Pro Ser Arg Leu Leu Arg Arg Ile Val His Met Glu Leu Gln Glu Ala Asp Gly Asp Cys His 30 Leu Gln Ala Val Val Leu His Leu Ala Arg Arg Ser Val Cys Val His 50 Pro Gln Asn Arg Ser Leu Ala Arg Trp Leu Glu Arg Gln Gly Lys Arg 60 Leu Gln Gly Thr Val Pro Ser Leu Asn Leu Val Leu Gln Lys Lys Met Tyr Ser Asn Pro Gln Gln Gln Asn 90 <210> 15 <211> 32 <212> DNA <213> Homo sapiens <400> 15 32 atctggcacc acaccttcta caatgagctg cg <210> 16 <211> 32 <212> DNA <213> Homo sapiens <400> 16 32 cgtcatactc ctgcttgctg atccacatct gc